

Figures

a.

Stresscopin 1

ATGACCAGGTGTCTGCTCTGCTGTTGCTGATGGTCTGATGTTGGGCAAGAGTC	51
M T R C A L L L M V L M L G R V	102
CTGGTTGTCCCAGTGACCCCTATCCCAACCTTCCAGCTCCGCCCTCAGAAT	
L V P V T P I P T F Q L R E Q N	
TCTCCCCAGACCACTCCCCGACCTCGGGCCTCAGAGAGCCCTCAGCTGCT	153
P Q T T P R P A A S E S P S A A	
CCCCACATGGCCGTGGGCTGCCAGAGCCA CTG CAG G C C A C C G C A C C C T	204
P T W P W A A D O S H C S P T R	
GGCTCGCGATTGCTCTATCGCTGGATGTCCCCATCGGCCTTGTGAGATC	255
TTACTGGAGCAAGCCGGGCCAGGGCTGCCAGGGAGCAGGCCACCACCAAC	306
GCCCGCATCCTGGCCGTGCGCCACTGCTGA	
G H C *	339

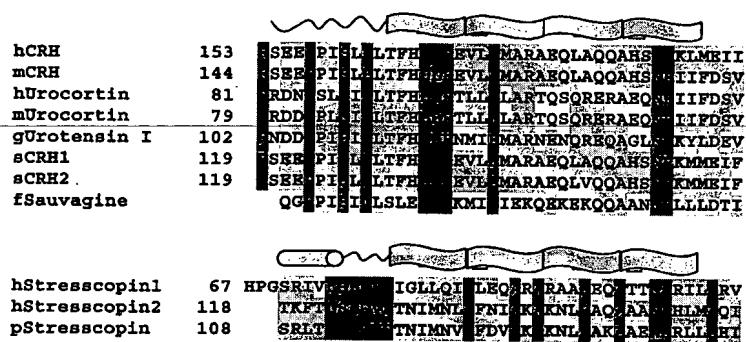
b.

Stresscopin 2

ATGCTGATGCCGCCACTTCTGCTGCTCCTGCTGCTGCTGGGGGGC	51
M L M P V H F L L L L L L L G G	102
CCCAGGACAGGCCCTCCCCACAAGTTCTACAAAGCCAAGCCCATCTTCAGC	
P R T G D P F H K F Y K A K P I F S	153
TGCCTAACACCGCCCTGCTGAGGCTGAGAAGGGCCAGTGGGAGGATGCA	
C L N T A L S E A E K G Q W E D A	
TCCCTGCTGAGCAAGAGGAGCTCCACTACCTGCGCAGAGACGCCCTCT	204
S L E S K R S F H Y L R S R D M S	
TGGGAGAGGAGGGAGGGAAAGAGAAAAAGACTTTCCCATCTCTGGG	255
S G E E E E G K E K K T F P I S G	
GCCAGGGGTGGAGCCGGAGGCACCCGTTACAGATACTGTCTCCAAGCACAG	306
A T R G G A G G T R M R Y V S O A Q	
CCCAGGGAAAGCCACGCCAGGACACAGCCAAGAGTCCCCACCGCACCAAG	357
P R A G K P R Q D T A K S P H R	
TTCACCTGTCCTCGACGCCACCATCATGAACCTCCTCTCAAC	408
ATCGCCAAGGCCAAGAACCTGCGTGCCAGGGCCGCAATGCCACCTG	459
ATGGCCCAAATTGGGAGGAAGAAGTAG	
G R R K *	486

Fig. 1a and 1b

C.



d.

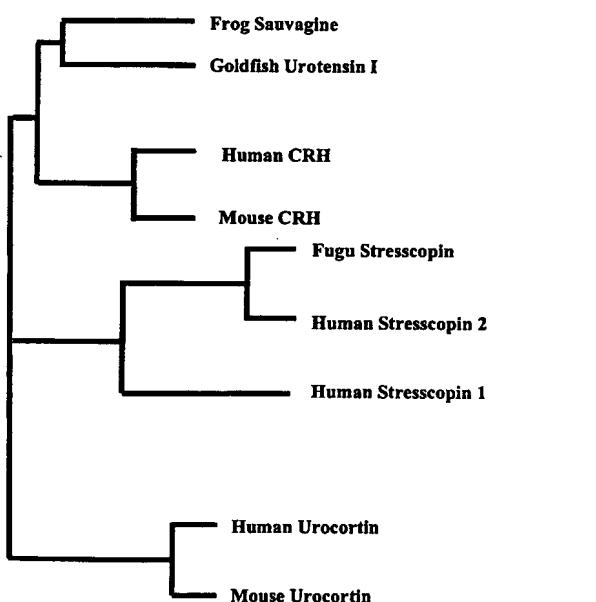


Fig. 1c and 1d

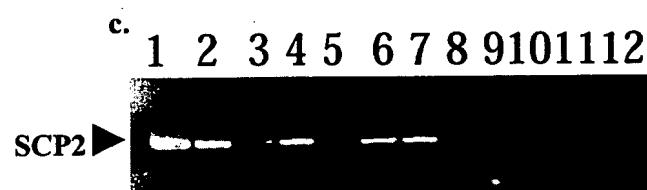
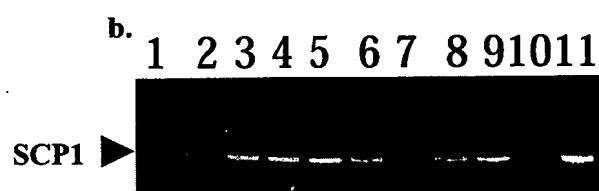
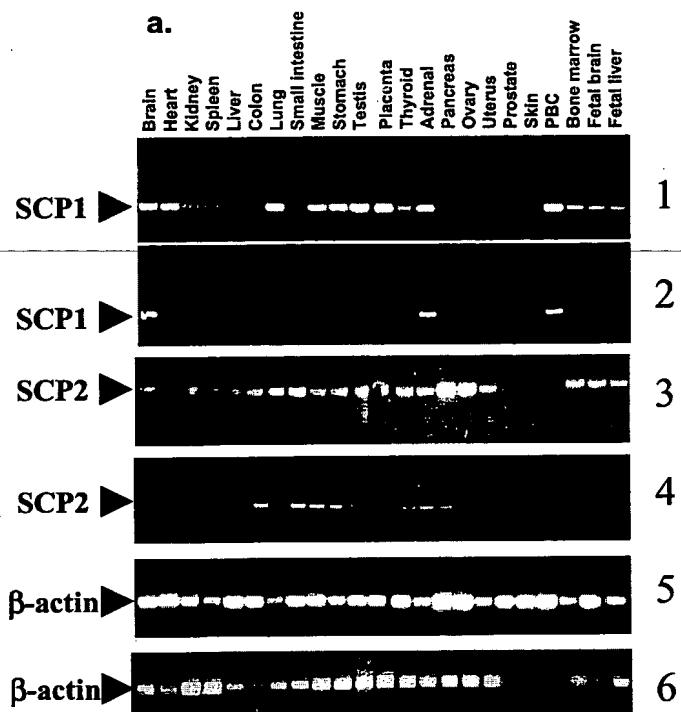
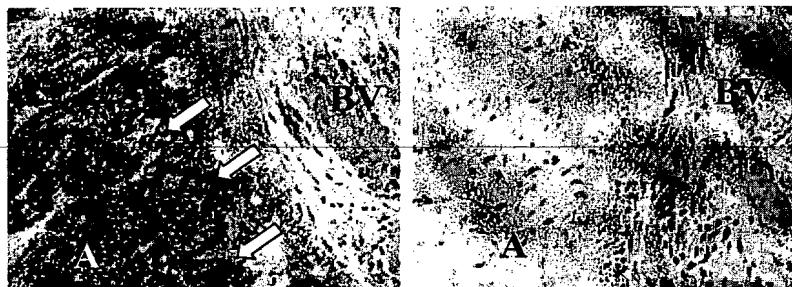
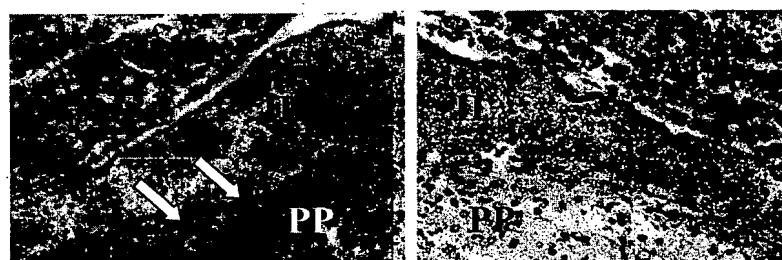


Fig. 2a-c

d.



e.



f.

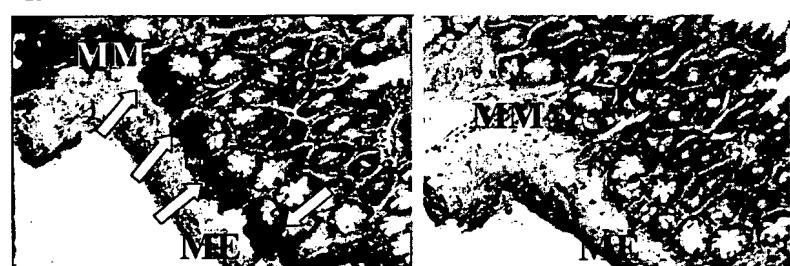


Fig. 2d-f

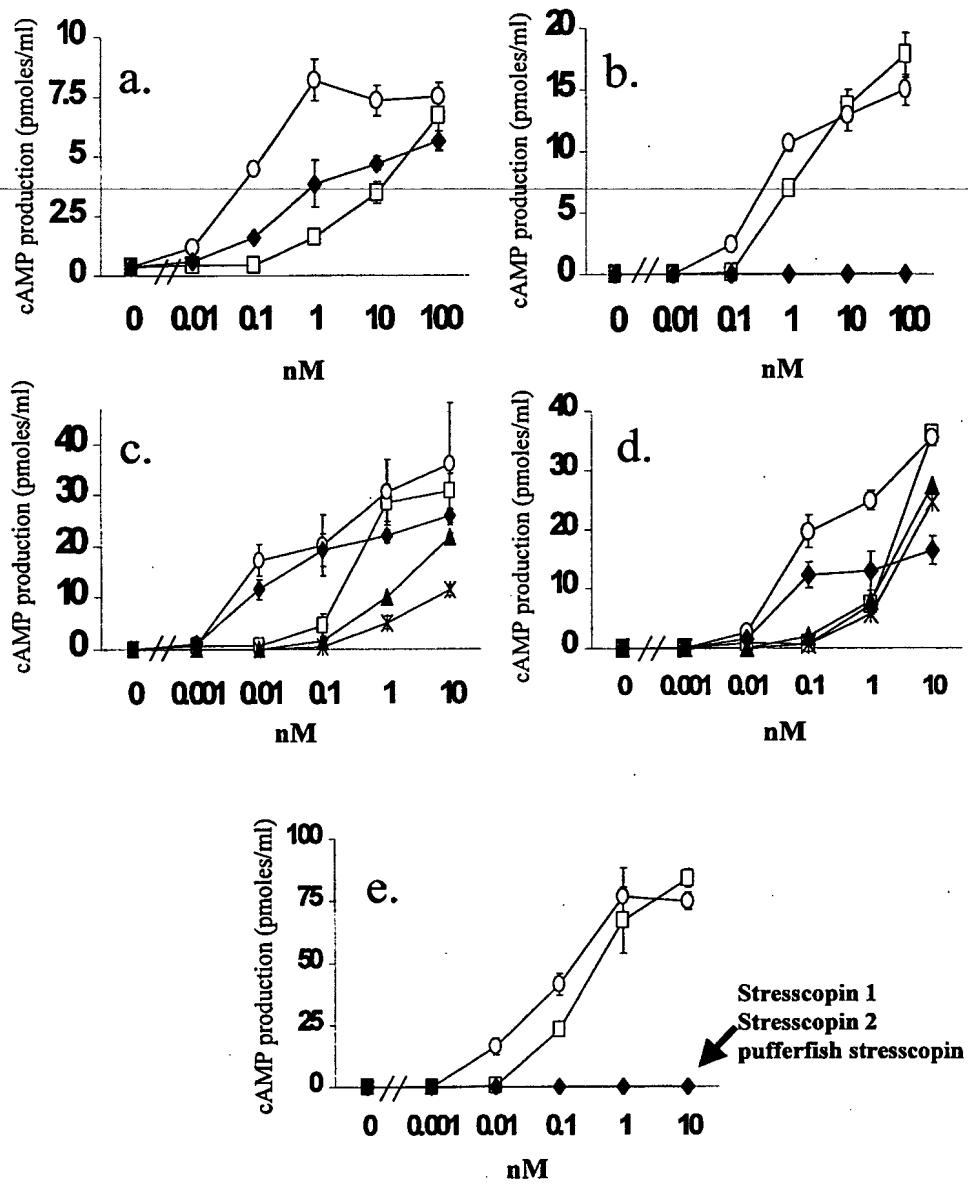


Fig. 3a-e

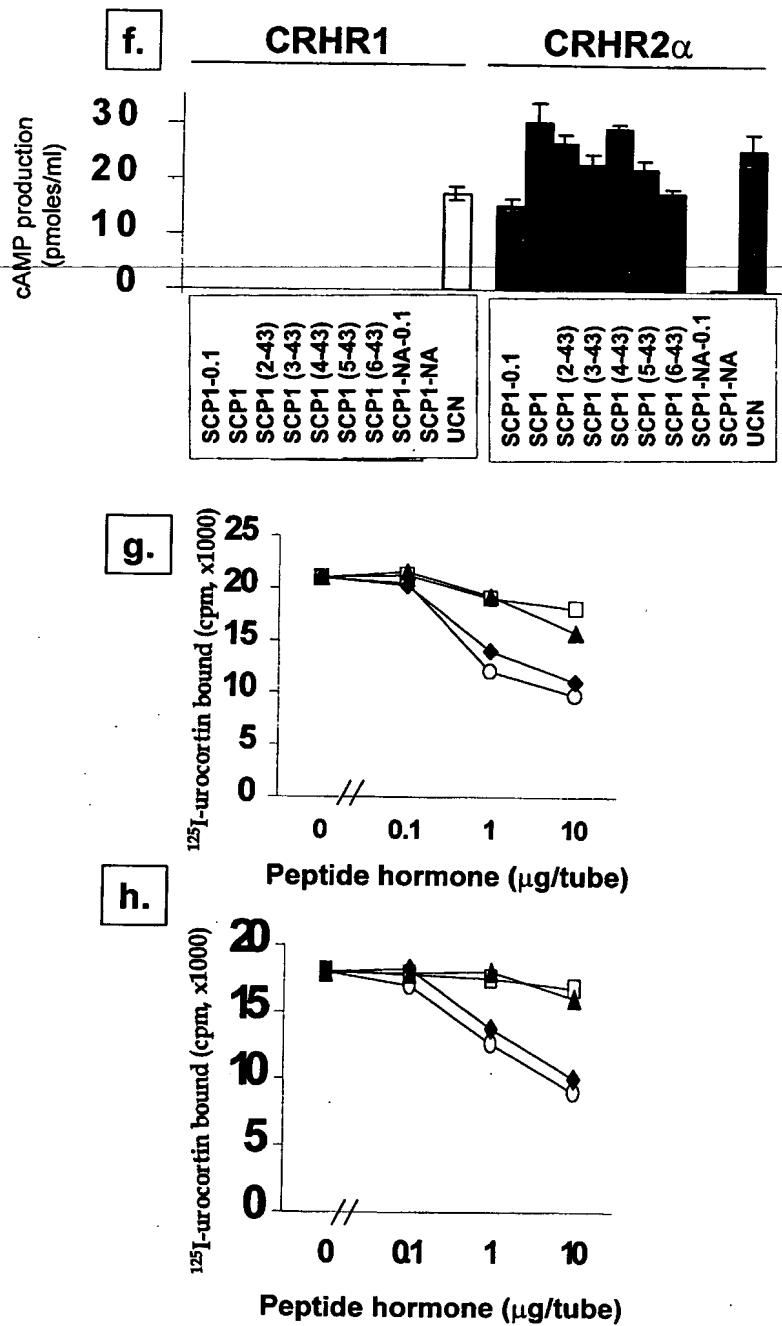


Fig. 3f-h

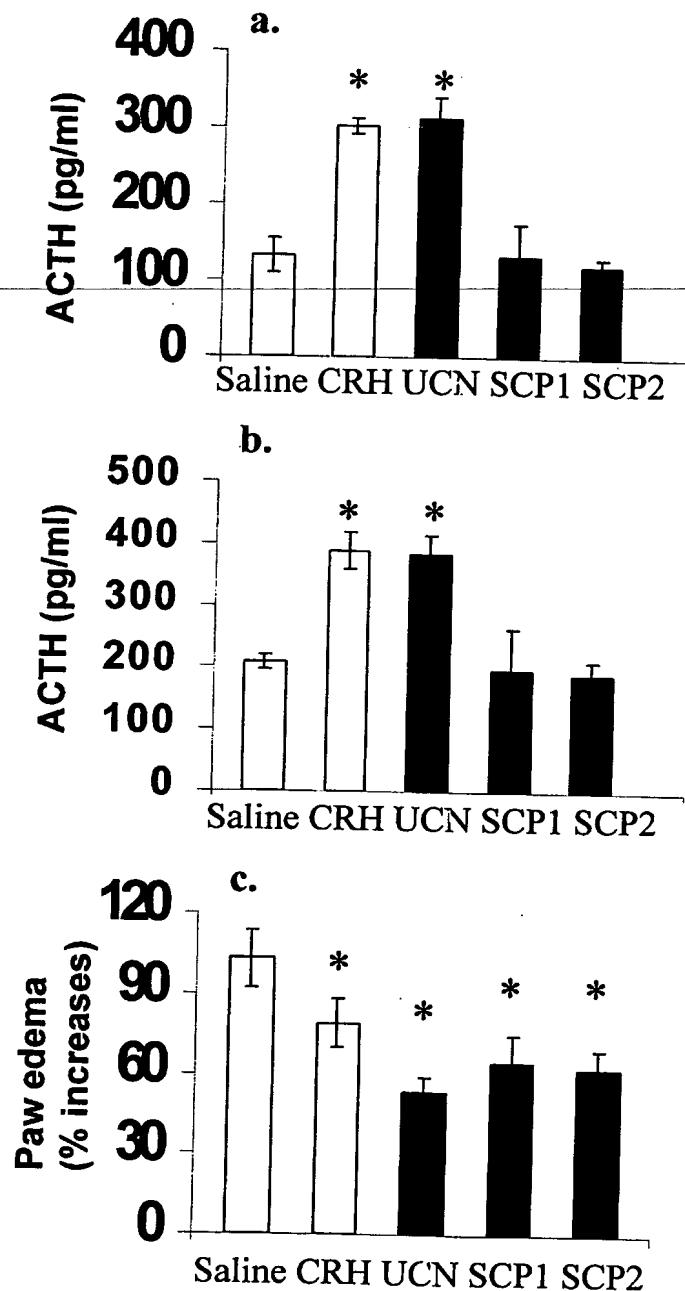
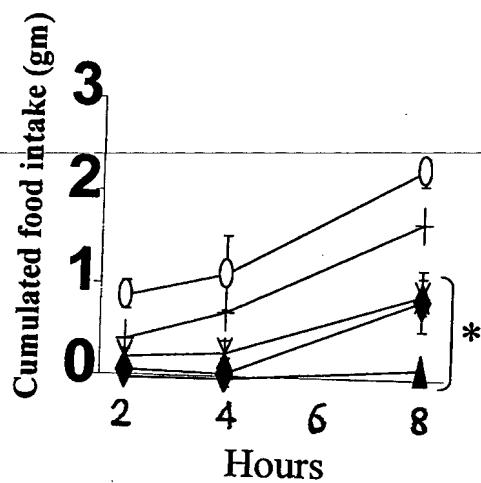
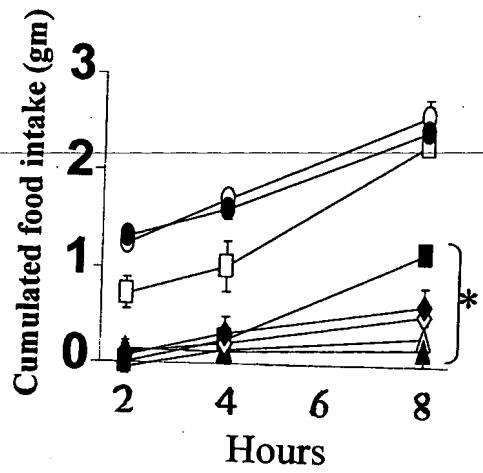


Fig. 4a-c

d.



e.

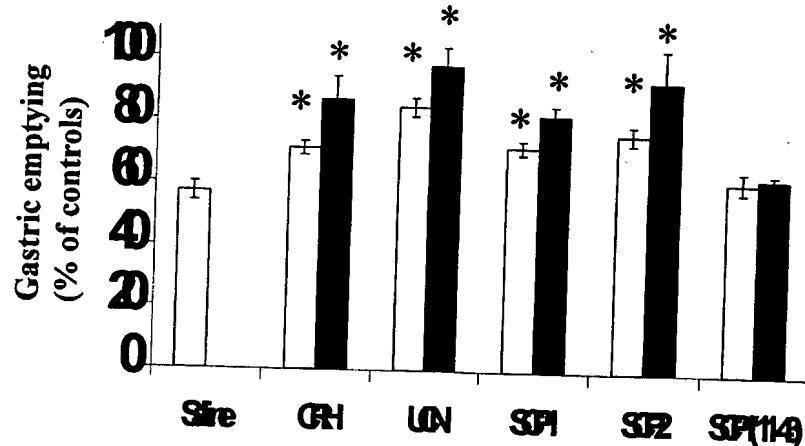


Fig. 4d-e